



[illegible][illegible]

[illegible]

Oct 4th 1948

DB 2592 CTAAAGTTTAAAGATTTTCTAGAAATAGCAAGAGCTGAAATTTGATTTTCTGCT 642
CY 643 ATTAT 702
DB 2642 ATGTGAGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 701
CY 703 CC 704
DB 2702 CC 2703

RESULT 12

US-09-021-287-2
Sequence 2, Application 35/09/46473
Patent No. 6297011
GENERAL INFORMATION:
APPLICANT: GreenSpan, Daniel S
APPLICANT: Takahara, Kazuhiko
APPLICANT: Hoffman, Guy G
TITLE OF INVENTION: Mammalian Toll-like protein
NUMBER OF SEQUENCES: 13
CORRESPONDENCE ADDRESS:
ADDRESSEE: Charles & Brady
STREET: 1 South Pinckney Street
CITY: Madison
STATE: WI
COUNTRY: US
ZIP: 53703
COMPUTER READABLE FORM:
MEDIUM TYPE: floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patented Release #1.0, Version #1.0
CURRENT APPLICATION DATA
APPLICATION NUMBER: 09/09/001,297
FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/09/001,297
FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Benson, Bennett J
REGISTRATION NUMBER: 37094
REFERENCE/PATENT NUMBER: 660266, 92819
TELECOMMUNICATION INFORMATION:
TELEPHONE: 608-251-5000
TELEFAX: 608-251-9166
INFORMATION FOR TEST NO. 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 4771 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
TOPOLOGY: linear
MOLECULE TYPE: DNA (genomic)
FEATURE:
NAME/REV: CDS
LOCATION: 611..3652
OTHER INFORMATION: (product: "murine mtl1 protein")
US-09-001 247 2

Query Match 2.47 Score 61.41 E-03 Length 4771
Best Local Similarity 50.841 Pctd No. 1.9e-057
Matches 123, Conservative 0, Mismatch 119, Indels 0, Gap 0
CY 443 TTTTAT 507
DB 1443 TTTTAT 1402
CY 643 ATTAT 702
DB 2642 ATGTGAGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 701

DB 583 CTAAAGTTTAAAGATTTTCTAGAAATAGCAAGAGCTGAAATTTGATTTTCTGCT 642
CY 643 ATTAT 702
DB 2642 ATGTGAGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 701
CY 703 CC 704
DB 2702 CC 2703

RESULT 13

US-09-021-240-473
Sequence 2, Application 35/09/46473
Patent No. 6297011
GENERAL INFORMATION:
APPLICANT: GreenSpan, Daniel S
APPLICANT: Takahara, Kazuhiko
APPLICANT: Hoffman, Guy G
TITLE OF INVENTION: Mammalian Toll-like protein
NUMBER OF SEQUENCES: 13
CORRESPONDENCE ADDRESS:
ADDRESSEE: Charles & Brady
STREET: 1 South Pinckney Street
CITY: Madison
STATE: WI
COUNTRY: US
ZIP: 53703
COMPUTER READABLE FORM:
MEDIUM TYPE: floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patented Release #1.0, Version #1.0
CURRENT APPLICATION DATA
APPLICATION NUMBER: 09/09/240,473
FILING DATE:
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Benson, Bennett J
REGISTRATION NUMBER: 37094
REFERENCE/PATENT NUMBER: 660266, 92819
TELECOMMUNICATION INFORMATION:
TELEPHONE: 608-251-5000
TELEFAX: 608-251-9166
INFORMATION FOR TEST NO. 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 4771 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
TOPOLOGY: linear
MOLECULE TYPE: DNA (genomic)
FEATURE:
NAME/REV: CDS
LOCATION: 611..3652
OTHER INFORMATION: (product: "murine mtl1 protein")
US-09-021 473 2

Query Match 2.47 Score 61.41 E-03 Length 4771
Best Local Similarity 50.841 Pctd No. 1.9e-057
Matches 123, Conservative 0, Mismatch 119, Indels 0, Gap 0
CY 443 TTTTAT 507
DB 1443 TTTTAT 1402
CY 643 ATTAT 702
DB 2642 ATGTGAGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 701



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62	1968-3			

REFSeq ID: A137891.1
 LOCUS: A137891.1
 DEFINITION: LOCUS_TAG: x1; Scores Nijmegen 31 Homo sapiens cDNA clone IMAGE:2794180 47, similar to TR-545460 (U937-545460) OTHERS: cDNA clone IMAGE:2794180
 TRANSMEMBRANE PROTEIN. 7; cDNA sequence.
 ACCESSION: A137891.1
 VERSION: A137891.1
 FEATURES: EST.
 SOURCE: human.
 ORGANISM: Homo sapiens
 Eukaryota; Metazoa; Chordata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominoidea; Homo.
 REFERENCE: 1 (bases 1 to 533)
 AUTHORS: NCI CCRP
 URL: <http://www.ncbi.nlm.nih.gov/ncbiinfo/>
 TITLE: National Cancer Institute, Cancer Genome Anatomy Project (NCAP), Tumor Gene Index
 JOURNAL: Published (1997)
 COMMENT: Contact: Robert Strauss, Ph.D.
 Email: roberts@ncbi.nih.gov
 This cDNA is available locally free through NCI's Cancer Gene
 (CAGE) program. Deposited by NCI for further information.
 Insert length: 1038. Std Error: 4369
 Seq primer: 740bp from Gibco
 High quality sequence stop: 44%.
 FEATURES: Location/Qualifiers
 1..533
 /organism="Homo sapiens"
 /db_xref="Gene:2794180"

[illegible]

